

Lubet

1644

Errors Corrected by the STIC Systems Branch

Serial Number: 08/737457

CRF Processing Date: 1/29/99
Edited by: Ar
Verified by: Ar (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☒ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

ENTERED

10 1/2

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/737,457DATE: 01/29/1999
TIME: 17:42:10

INPUT SET: S30407.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

Does Not Comply
Corrected Diskette Needed

1 SEQUENCE LISTING

3 (1) General Information:

4 (i) APPLICANT:

- > 5 (A) NAME: BIOVATION LIMITED
--> 6 (B) STREET: Crombie Lodge, Aberdeen Science Park, Balgownie Drive
--> 7 (C) CITY: Aberdeen
--> 8 (E) COUNTRY: United Kingdom
--> 9 (F) POSTAL CODE (ZIP): AB22 8GU
10

11
12 (ii) TITLE OF INVENTION: Improvements in or Relating to Peptide
13 Delivery
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15 (iii) NUMBER OF SEQUENCES: 6
16

17 (iv) COMPUTER READABLE FORM:

- 18 (A) MEDIUM TYPE: Floppy disk
19 (B) COMPUTER: IBM PC compatible
20 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
21 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
22

23 (vi) curr. APP. DATA: *add leading*

24 (2) INFORMATION FOR SEQ ID NO: 1:

25 (i) SEQUENCE CHARACTERISTICS:

- 26 (A) LENGTH: 36 base pairs
27 (B) TYPE: nucleic acid
28 (C) STRANDEDNESS: single
29 (D) TOPOLOGY: linear
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36 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
37

38 GCAGCCGCGA AGTATATCTG CAACTCATCC TGCATG
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36

40 (2) INFORMATION FOR SEQ ID NO: 2:

41 (i) SEQUENCE CHARACTERISTICS:

- 42 (A) LENGTH: 36 base pairs
43 (B) TYPE: nucleic acid
44 (C) STRANDEDNESS: single
45 (D) TOPOLOGY: linear
46

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/737,457DATE: 01/29/1999
TIME: 17:42:10

INPUT SET: S30407.raw

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

CATGCAGGAT GAGTTGCAGA TATACTTCGC GGCTGC

36

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

GCAGCCGCGG GAATACTAGG GTTCGTATTT ACACTA

36

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

TAGTGTAAT ACGAACCTA GTATTCCCGC GGCTGC

36

(2) INFORMATION FOR SEQ ID NO: 5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 42 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/737,457DATE: 01/29/1999
TIME: 17:42:10

INPUT SET: S30407.raw

100 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
101
102 GGCCGCTACG GACGAAAGAA GAGGCGTCAA CGACGCAGAC CA 42
103
104 (2) INFORMATION FOR SEQ ID NO: 6:
105
106 (i) SEQUENCE CHARACTERISTICS:
107 (A) LENGTH: 42 base pairs
108 (B) TYPE: nucleic acid
109 (C) STRANDEDNESS: single
110 (D) TOPOLOGY: linear
111
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116 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
117
118 TGGTCTGCGT CGTTGACGCC TCTTCTTTTCG TCCGTAGCGG CC 42
119

PAGE: 1

SEQUENCE MISSING ITEM REPORT
PATENT APPLICATION US/08/737,457

DATE: 01/29/1999
TIME: 17:42:11

INPUT SET: S30407.raw

ADDRESSEE
STREET
CITY
STATE
COUNTRY
ZIP
CORRESPONDENCE ADDRESS
APPLICATION NUMBER
FILING DATE
CLASSIFICATION
CURRENT APPLICATION DATA
APPLICATION NUMBER
FILING DATE
PRIOR APPLICATION DATA